



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 166874

To: **Ruixiang Li**
Location: **REM-4D75&4C70**
Art Unit: **1646**
Friday, November 05, 2004

Case Serial Number: **09/801944**

From: **Beverly Shears**
Location: **Remsen Bldg.**
RM 1A54
Phone: **571-272-2528**

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Search Notes

Protein Sequence Searches - 10/8/04

All of the sequence databases on the ABSS have been updated. A change has occurred in the protein databases.

- Two protein databases, SPTREMBL and SwissProt, are now produced as a single, merged database called UniProt.
- Results from UniProt have the file extension **.rup**.
- Sequences in UniProt are identified by the same ID that had been used in SPTREMBL or SwissProt.
- In instances where the database curators have determined that an SPTREMBL record and a SwissProt record represent the same sequence, the two records have been merged into one. Both IDs are present in the record. Any differences found between the two sequences are recorded in the FT (feature table) fields.

If you have any questions regarding these changes or your results, please contact any STIC searcher.

From: Li, Ruixiang
Sent: Tuesday, November 02, 2004 12:13 PM
To: STIC-Biotech/ChemLib
Subject: Sequence search of Application No.09/801,944

Please do a standard search on:

SEQ ID NO: 268 against both commercial and interference amino acid databases.

321 PA

CRF
Thank you very much!

Ruixiang Li
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REM 4D75
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STAFF USE ONLY

Searcher: _____
Searcher Phone: 2-
Date Searcher Picked up: _____
Date Completed: _____
Searcher Prep/Rev. Time: _____
Online Time: _____

Type of Search
NA Sequence: # _____
AA Sequence :# _____
Structure: # _____
Bibliographic: _____
Litigation: _____
Patent Family: _____
Other: _____

Vendors and cost where applicable
STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other(Specify): _____

Date completed: _____
Searcher: Beverly e 2528
Terminal time: _____
Elapsed time: _____
CPU time: _____
Total time: _____
Number of Searches: _____
Number of Databases: _____

Search Site
____ STIC
____ CM-1
____ Pre-S

Type of Search
____ N.A. Sequence
____ A.A. Sequence
____ Structure
____ Bibliographic

Vendors
____ IG
____ STN
____ Dialog
____ APS
____ Geninfo
____ SDC
____ DARC/Questel
____ Other CRF